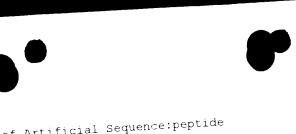
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    Met Hie Al & Gly
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tgotototaa gggtotaoto	
KD1W+ 3	
KU11 - 15	
ROID - DIA	
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<pre>&220</pre>	
<pre>2.13 · Description of Artiffer</pre>	
. 4005 3	15
ctctaaggta aatat	
$\pm i \cos 4$	
311 · 16	
(212 · DNA	
213 Simian virus 40	
200 - Sequence:peptide	
2000 Description of Artificial Sequence:peptide	
. 400 - 4	16
tgtattttag attcaa	
. 210 - 5	
3.211 × 19	
CION DNA	
.213% Simian virus 40	
.220 Description of Artificial Sequence:peptide	
. 223 - Description of Alteriable	
.400 + 5	19
tgctctctaa ggtaaatat	
; 1× €	
00110 19	
COLOR DNA	
213> Simian virus 40	
200 · 203 · Description of Artificial Sequence:peptide	
23. Description of Alexander	
400 > 6	19
tytattttag ggtctactc	
.,∠10> 7	

<pre><:11 + 19</pre>	
KO10 PNA	
RECTION Bacteriophage T7	
*310 · sequence:peptide	
<pre>% Dubble</pre>	
$4 \hat{n} \hat{n} + 7$	19
_{ய்று} நம்பும் புக்க gguaaauau	
- <u> </u>	
.211 - 19	
- 011 - 10 - 010 - RNA	
213 Bacteriophage T7	
no de la companya de	
• 2000 ·	
	19
400 + 8	
uguauuuuag ggucuacuc	
- 2101 9	
- 211 + 20	
C12 FNA	
- 213 - Bacteriophage T7	
220 · Cartificial Sequence:peptide	
- 220 223 - Description of Artificial Sequence:peptide	
. 400× 9	20
ugoucucuaa gggucuacuc	
.0105-10	
- 2115 49	
anion dna	
:213 · Bacteriophage T7	
(
003 - Description of Artificial Sequence:peptide	
- 400 - 10	49
-400 - 10 ggaattogto togagototg atcaccacca tggacacgat taacatogo	
. <u>210</u> - 11	
111 · 55	
212 · DNA	
(21) Basteriophage T7	
(21) - 000001-1	



<pre><:000</pre>	
$< 4 \mathrm{min} \cdot 11$ gurtlagttigg tetegtetet tittiggagg agtgiegtie tiagegatgi taate	55
<pre>cole + 10 col = 46 col = DNA col = Bacteriophage T7</pre>	
.220 · Cescription of Artificial Sequence:peptide	
400×12 agaattogto toggagaaag gtaaaattot otgacatoga actggo	46
2108 13 -211 - 33 -212 DNA -213 Bacteriophage T7	
220 Pescription of Artificial Sequence:peptide	
.400 × 13 gadtagtggt otocoottag agagoatgto ago	33
00100 14 0011	
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.,tb/14 _{Ny} aatteggt otogggtota otoggtggog agg	33
.210:-15 .111 · 27 .212 DNA .113 · Bacteriophage T7	
- 200 · . .203 · Description of Artificial Sequence:peptide	

4

(400 ≥ 15





	21
gantallingt tangegaang caaagto	
<510 ± 16	
- 211 · 36	
and the principal of the control of	
Easteriophage T7	
- 220	
.400 · 16	36
(400 · 16 ggaattogto totaaggtaa atataaaatt tttaag	
210 - 17	
211 × 40	
TO TONA	
- 213 - Bacteriophage T7	
- 220 · C. Patificial Sequence:peptide	
220 Description of Artificial Sequence:peptide	
φήφ - 17	40
400×17 gactagtogt etetgaceet aaaatacaca aacaattaga	
. <u>210</u> - 18	
- 311 - 92	
. 112 · DNA	
Bacteriophage T7	
2208 2238 Description of Artificial Sequence:peptide	
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ggaattegte tegagetetg accaactag te	
ggaattegte tegageeers eteetesaaa aaagagaega gaccaactag te	
1:(> 19	
. <u>211</u> 2-92	
CLES DNA	
113. Bacteriophage T7	
400 > 19 gaztagttig getegtetet tittiggagg aggggegtte tiagegatgt	taatcgtgtc 60
gactagttigg gotogtotol tectoggass	
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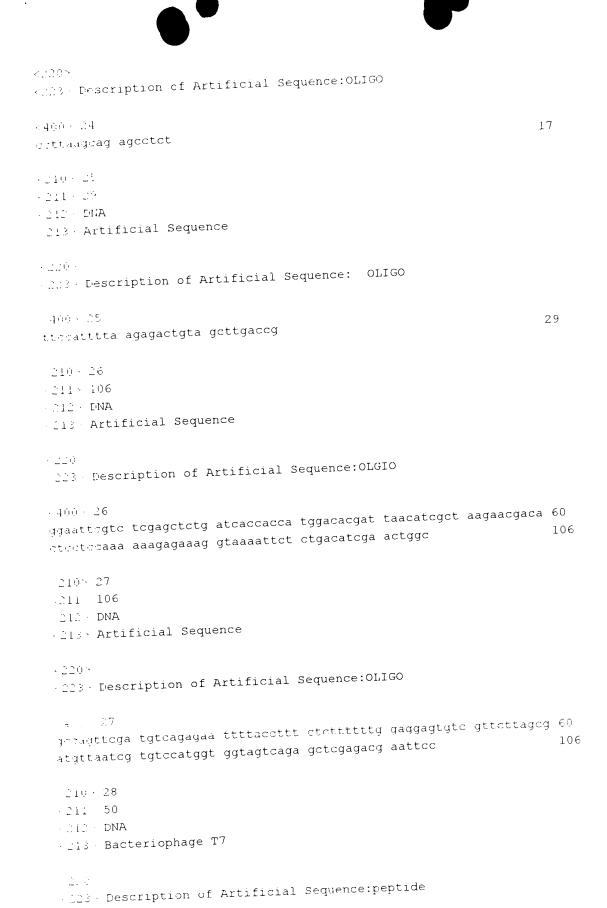




33

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.220% .228 Description of Artificial Sequence:peptide
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.210% 22 .211
<pre>Ship Ship Ship Ship Ship Ship Ship Ship</pre>
.400> 32 ggaattegte teg
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- .213 Artificial Sequence .223 Description of Artificial Sequence: OLIGO . 400 - 23 qagaaaggta aaattototg acatogaact ggo
 - .210 24 - 211 - 17 212 DNA







$< 400 \pm 10^3$ atggassaga ttaacatege taagaacgae ttetetgaca tegaactgge	50
210 - 29	
- < 211 - 50	
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.400 / 29 godagttoga tgtcagagaa gtogttotta gogatgttaa togtgtocat	50
gja - 30	
- 311 - 77	
NIN THIA	
-213 - Human immunodeficiency virus	
<pre>0220 * 0223 * Description of Artificial Sequence:peptide</pre>	
CDL3 - Description	
. 100 × 30	attc 60
(400 × 30) auggasacga ttaacatogo taagaacgao actootocaa aaaagagaaa ggtaaa	77
totgacatog aactggc	
1.00 6 4 M 2 M 2 M 2 M 2 M 2 M 2 M 2 M 2 M 2 M	
. 210% 31	
. 211> 77	
- 212 - DNA	
013 · Bacteriophage T7	
400% 31 geragttega tgteagagaa ttttacettt etettttttg gaggagtgte gttet atgttaateg tgteeat	tagcg 60 77
210 - 32	
.011 - 69	
AUG. DIA	
- :::3 - Human immunodeficiency virus	
<pre>- 2000 2003 - Description of Artificial Sequence:peptide</pre>	
400 - 32	tgctta 60
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agootcaag	
- 010 + 33	
- 111 - 69	
L11 DNA	
Control Diffe	





<pre><ii13 human="" immunodeficiency="" pre="" virus<=""></ii13></pre>
(200)
CLIDUS CLIDUS Description of Artificial Sequence:peptide
 (4)00 - 33 gatrottyag gottaagcag tgggttooct agttagccag agageteeca ggctcagate 60
tggt staat
- <u>210</u> - 34
> 211 + 61
- NIN - ENA - NIS - Human immunodeficiency virus
- C20 ·
- 220 - Description of Artificial Sequence:peptide
400 × 34
400×34 gatoacetta ggototoota tggoaggaag aagoggagao agogaogaag acotootoaa 60
a a constant of the constant o
. <u>210 - 35</u>
£11 × 61
212 - DNA
213 · Human immunodeficiency virus
220 -
DD3 Description of Artificial Sequence:peptide
. 4005 35
- 400 > 35 gateettgag gaggtetteg tegetgtete egettettee tgeeatagga gageetaagg 60
t
- 210 - 36
. 111 - 62
212 · DNA
113 · Human immunodeficiency virus
320
229 - 223 - Description of Artificial Sequence:peptide
400 · 36
$400 \cdot 36$ $_{ m patch}$ atagtg aatagagtta ggcagggata otoaccatta togtttoaga ocoacctooc 6
ag

WILL + DNA

210 - 37 -211 - 62





28

<.:13 -	Facteriophage	T7
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gatortggga ggtgggtotg aaacgataat ggtgagtato ootgootaac totattoact 60

- v 210 × 38
- $\times 211 \times 30$
- 213 + BHA
- []] Artificial Sequence
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- Lib Description of Artificial Sequence: oligo

30 .400 - 38 matctagage taacaaagee egaaaggaag

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- Z13 ENA
- -313 Artificial Sequence
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ttotgoagat atagttooto otttoago

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- 212 DNA
- -113 Artificial Sequence
- 020 -
- Description of Artificial Sequence:oligo
- $..400 \leftarrow 40$

togagedatg gettaaggat degtaegtee ggagetageg ggedeatega tactagttaa 60 atgragatct

- ...10 41
- 11 70
- ._!_ DNA
- Artificial Sequence





union de la companya	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
$<\!400\cdot41$ cragagatet geatttaact agtategatg ggeoogetag eteoggaegt aeggateett	60
<pre><400 - 41 **The grant taact agtategatg ggcccgctag etccggacgt acggacos."</pre>	70
ctagadatic godos	
aaq राज्यं पुर्व	
7 (210 × 42)	
.311 · 0#	
CALC - ENA Sequence	
3213 Artificial Sequence	
-1:40	
220 · Description of Artificial Sequence:oligo	
. [23] - Description	
	29
-400 - 40 catgaaatta attogaotoa otataogga	
_{Datiq} aaatta acceguduu	
1.010 ± 43	
(211) 29	
212 LMA	
213 Artificial Sequence	
li mo	
- 320	
. Dog. Description	
12	29
.400> 43 tttaattaag otgagtgata tgoototag	
tttaattaay cogus s	
. <u>2103 44</u>	
- D11 - 72	
CD10 DNA CD13 Artificial Sequence	
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- 220 ·	
$_{ m cut}$ $_{ m$	ctcag 60
44V 444 gaggettaag cagtgggtte ectagttage edgags	72
gatinggaco gess	
arctypicta at	
212. 45	
. 210 + 45	
- 111 + 72	
212 · DNA 213 · Artificial Sequence	
Size Artificana a r	
11.70	
0.2235 Description of Artificial Sequence:oligo	
.2230 Descripción	





k4000 45 Goggattaga coagatotga gootgggago tototggota actagggaad coactgotta 60 72 agcotcaato og
₹310 × 46
- DII - 66
AM2 AM2
213 · Artificial Sequence
<pre>.020</pre>
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.210× 47
CALAS TRIA
: 313 · Artificial Sequence
<pre>.2200 .223. Description of Artificial Sequence:oligo</pre>
400 imes47 coggaeetta ggeteteeta tggeaggaag aageggagae agegaegaag aceteeteaa 60
ggtccg
KC108 48
. <u>211</u> ~ 65
1115 DNA
C13 Artificial Sequence
- 2005 2003 Description of Artificial Sequence:oligo
4000 48 gatucggatg ggaggtgggt otgaaacgat aatggtgagt atcootgoot aactotatto 60
gatucggacg ggugg-ssss
actat
∴10° 49
_11 · 65
312 - DNA
213 Artificial Sequence
nno:
220) 0.223> Description of Artificial Sequence:oligo





$_{\rm C400^{5}}$ 49 coggata etcaccatta tegitteaga eccacetece 60 coggatagig aatagagita ggeagggata etcaccatta tegitteaga eccacetece 60 atong
*2105 50 *211 * 67 *212 * DNA *213 * Artificial Sequence
.220 > .223 > Description of Artificial Sequence:oligo
400> 50 Jatcagcatg cetgeaggte gaetetagae eegggtaeeg agetegeeet atagtgagte 60 67 gtattat
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.220 · Description of Artificial Sequence:peptide
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